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A THESIS FOR THE DEGREE OF MASTER

Fibroblast growth factor-1 as a mediator of paracrine  
effects of canine adipose tissue-derived  
mesenchymal stem cells on *in vitro*-induced insulin  
resistance models

*In vitro* 인슐린 저항성 모델에 대한 개  
지방유래중간엽줄기세포 paracrine 효과의 매개체  
fibroblast growth factor-1

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김현진

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이 논문을 수의학석사학위논문으로 제출함.

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**Fibroblast growth factor-1 as a mediator of paracrine effects of canine  
adipose tissue-derived mesenchymal stem cells on *in vitro*-induced  
insulin resistance models**

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## **Abstract**

In the field of diabetes research, many studies on cell therapy have been conducted using mesenchymal stem cells. This research was intended to shed light on the influence of canine adipose-tissue-derived mesenchymal stem cell conditioned medium (cAT-MSC CM) on *in vitro* insulin resistance models that were induced in differentiated 3T3-L1 adipocytes and the possible mechanisms involved in the phenomenon. Gene expression levels of insulin receptor substrate-1 (IRS-1) and glucose transporter type 4 (GLUT4) were used as indicators of insulin resistance. Relative protein expression levels of IRS-1 and

GLUT4 were augmented in the cAT-MSC CM treatment group compared to insulin resistance model controls, indicating beneficial effects of cAT-MSC to DM, probably by actions of secreting factors. With reference to previous studies on fibroblast growth factor-1 (FGF1), it was proposed as a contributing factor to the mechanism of action. Anti-FGF1 neutralizing antibody was added to the CM-treated insulin resistance models. As a result, significantly diminished protein levels of IRS-1 and GLUT4 were observed, supporting our assumption. Accordingly, this study advocated the potential of MSC CM as an alternative insulin sensitizer and discovered a signalling factor associated with the paracrine effects of cAT-MSC.

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**Key words:** fibroblast growth factor-1; canine adipose tissue-derived mesenchymal stem cells; conditioned medium; diabetes mellitus; insulin resistance; in vitro; 3T3-L1

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# 1. Introduction

Diabetes mellitus (DM) is an important endocrine disease that accounts for a serious proportion of small animal medicine. In dogs, most DM patients are known to be type 1 DM (T1DM), which is usually well managed by exogenous insulin supplements. However, canine DM with insulin resistance, ‘poorly controlled T1DM’, not only needs expensive treatment, but also has poor prognosis. In human medicine, several reports advocate that concurrent diseases such as obesity and inflammatory diseases are associated with insulin resistance in DM patients [7, 9, 38, 48]. A study in dogs also asserts the obesity is a cause of insulin resistance [13, 14]. However, the exact pathophysiology of insulin resistance development is not fully understood. Although insulin sensitizers could be an option for the treatment of uncontrolled T1DM, their possible adverse effects such as weight gain, bone loss, and congestive heart failure encourage further effort to develop new drugs for insulin resistance [28].

Mesenchymal stem cells (MSCs) are multipotent stromal cells that have immunomodulatory and regenerative effects. Because they are relatively free from ethical issues, their therapeutic uses for various diseases including DM have been studied globally. Many studies have shown that they have advantageous effects in *in vivo* experiments using diabetic rodent models [3,



6, 11, 16, 23, 25]. Until now, the known mechanisms of MSC actions when applied to DM are as follows: to differentiate directly into insulin-producing cells (IPCs), to regulate the immune system, and to secrete beneficial cytokines and growth factors [50]. In particular, the paracrine effects of MSCs are thought to enhance insulin sensitivity [40].

Fibroblast growth factor-1 (FGF1) which is a member of the FGF family has been known to play crucial role in glucose homeostasis [31]. Perry et al. reported that FGF1 and FGF19 improved glucose metabolism via down regulation of the hypothalamic-pituitary-adrenal axis [33]. It has been documented that exogenous recombinant FGF1 (rFGF1) improved insulin sensitivity as well as normalized blood glucose levels in diabetic mice models [42]. Recently, FGF1 is getting attention as a leading candidate of novel insulin sensitizer.

This study was designed to investigate the effects of canine adipose tissue-derived MSC-conditioned medium (cAT-MSC CM) on an *in vitro* induced insulin resistance model. To explore this, gene expression of markers related to glucose uptake were evaluated and the specific effective factors were discovered.

## 2. Material and Methods

### *2.1. Cell culture and characterization*

Adipose tissue was obtained from a healthy dog (1-year-old) during ovariectomy at the Seoul National University Veterinary Medicine Teaching Hospital (SNU VMTH). The owner was provided an informed, written consent for research use. The procedure was also approved by the Institutional Animal Care and Use Committee (IACUC) of SNU and the protocol was performed in accordance with the approved guidelines. Canine adipose tissue-derived mesenchymal stem cells (cAT-MSCs) were isolated and cultured as previously described [26]. Before their use in this study, cells were characterized by their ability to express several stem cell markers using flow cytometry. The following antibodies were used: cluster of differentiation (CD) 29-Fluorescein isothiocyanate (FITC), CD31-FITC, CD34-phycoerythrin (PE), CD73-PE (BD Biosciences, Franklin Lakes, NJ, USA), CD44-FITC, CD45-FITC, and CD90-allophycocyanin (APC) (eBiosciences, San Diego, CA, USA)-conjugated antibodies. Cells were analysed by a BD FACS Aria II system (BD Biosciences). Cellular differentiation was confirmed using PRIME-XV® Chondrogenic Differentiation Xeno-Free Serum-Free Medium (XSFM), PRIME-XV® Osteogenic Differentiation Serum-Free Medium (SFM), and PRIME-XV®

Adipogenic Differentiation SFM (all from Irvine Scientific, Santa Ana, CA, USA) according to the manufacturer's instructions followed by Alcian Blue staining, Alizarin Red staining, and Oil Red O staining, respectively.

## ***2.2. In vitro cellular insulin resistance models***

The *in vitro* induced insulin resistance model was developed in differentiated 3T3-L1 adipocytes. Murine 3T3-L1 cells were purchased from the Korean Cell Line Bank (Seoul, Korea). 3T3-L1 preadipocytes were differentiated using 3T3-L1 Differentiation Kit (Sigma-Aldrich) and all the differentiation procedures performed following the manufacturer's instructions. Once the proportion of differentiated cells reached 70%, induction of insulin resistance was initiated. Cells were washed with phosphate-buffered saline (PBS; PAN Biotech, Aidenbach, Germany) and changed to serum-free Dulbecco's modified Eagle's medium: Nutrient Mixture F-12 (1:1) (DMEM / F12 (1:1); PAN Biotech). Insulin resistance was induced with treatment with both 40 ng / mL of tumour necrosis factor- $\alpha$  (TNF- $\alpha$ ; PeproTech, Rocky Hill, NJ, USA) and 1% oxygen for 24 h. For hypoxic incubation, cells were placed in a hypoxic incubator (ViVAGEN, Sungnam, Korea) with the conditions of 1% O<sub>2</sub> at 37 °C.

### ***2.3. Preparation of cAT-MSC-conditioned medium (CM), and anti-fibroblast growth factor-1 (FGF1) antibody treatment***

cAT-MSCs ( $3 \times 10^5$  cells / well) were seeded in 6-well plates and cultured in 3 mL of DMEM medium containing 10% fetal bovine serum (FBS; PAN Biotech) for 3 days, yielding conditioned medium. After 3 days, conditioned medium was harvested and centrifuged at 850 rpm for 3 min to remove cellular debris. After centrifugation, the supernatant was transferred to a conical tube and stored at -80 °C until use. Anti-FGF1 neutralizing antibody (Abcam, Cambridge, MA, USA) was added to final concentrations of 0.9 µg / ml of culture medium.

### ***2.4. RNA extraction, cDNA synthesis, and quantitative real-time PCR***

Total RNAs from all cell groups were extracted using the Easy-BLUE Total RNA Extraction kit (Intron Biotechnology, Sungnam, Korea). The concentration and purity of RNA samples were measured by a spectrophotometer (Implen, Munich, Germany). First strand cDNA was synthesized using LaboPass M-MuLV Reverse Transcriptase (Cosmo

Genetech, Seoul, Korea) according to the manufacturer's instructions. Quantitative real-time (qRT)-PCR analysis was performed in duplicate in 96-well plates with StepOne Real-Time PCR System (Applied Biosystems, Foster City, CA, USA) using 10  $\mu$ L of AMPIGENE qPCR Green Mix Hi-ROX with SYBR Green dye (Enzo Life Sciences, Farmingdale, NY, USA) and 400 nM forward and reverse primers (Cosmo Genetech). The cycling conditions were as follows: polymerase activation for 2 minutes at 95 °C, followed by 40 cycles of denaturation for 5 seconds at 95 °C and annealing for 25 seconds at 60 °C. Expression levels were normalized to that of glyceraldehyde 3-phosphate dehydrogenase (GAPDH).

## ***2.5. Western blot analysis***

Total proteins were extracted from all cell groups using PRO-PREP Protein Extraction Solution (Intron Biotechnology) according to the manufacturer's instructions. The concentrations of the protein samples were measured using the Bio-Rad DC Protein Assay Kit (Bio-Rad, Hercules, CA, USA). The proteins were separated by sodium dodecyl sulphate-polyacrylamide gel electrophoresis and transferred to polyvinylidene difluoride membranes (Millipore, Billerica, MA, USA). The membranes were blocked by 5% non-fat dry milk in Tris-buffered saline containing 0.1%

Tween 20 and incubated with primary antibodies against insulin receptor substrate 1 (IRS-1, 1:500; Abcam) and glucose transporter type 4 (GLUT4, 1:500; Santa Cruz Biotechnology, Santa Cruz, CA, USA) at 4 °C overnight. The membranes were incubated with secondary antibodies at room temperature for 1 h. The immunoreactive bands were visualised using enhanced chemiluminescence (Advansta, Menlo Park, CA, USA) and normalised to  $\beta$ -actin levels (Santa Cruz Biotechnology).

## ***2.6. Statistical analysis***

Data are shown as mean  $\pm$  standard deviation. Statistical comparisons between groups were made with use of one-way ANOVA and an unpaired Student's t test using the GraphPad Prism v.6.01 software (GraphPad Inc., La Jolla, CA, USA). P value of  $< 0.05$  was considered statistically significant.

## 3. Results

### ***3.1. Characterization of cAT-MSCs***

Cells obtained from canine adipose tissue were characterized by their ability to express stem cell markers and to differentiate toward adipogenic, osteogenic, and chondrogenic lineages when cultured in media containing lineage-specific factors. The known MSC markers such as CD29, CD73, CD44, and CD90 were highly expressed by the cells. Negative markers such as CD31, CD34, and CD45 were not expressed (Figure. 1a). The multi-lineage plasticity of cAT-MSCs was confirmed by specific staining methods: Oil Red O staining, Alizarin Red S staining, and Alcian Blue staining, respectively (Figure. 1b).

### ***3.2. Differentiation into adipocytes and development of an insulin resistance model in 3T3-L1 adipocytes***

All processes of differentiation and induction of insulin resistance in the 3T3-L1 are depicted schematically in Figure 2a. During differentiation, 3T3-L1 preadipocytes transformed from a fibroblast-like appearance to an adipocyte-like appearance and cytoplasmic lipid droplet accumulation

became remarkable (Figure. 2b-d). After over 70% of 3T3-L1 cells differentiated, cells were treated with TNF- $\alpha$  and incubated in hypoxic conditions for 24 h to induce insulin resistance. The mRNA expression levels of IRS-1 and GLUT4 were compared as markers of insulin resistance. As a result, both IRS-1 and GLUT4 mRNA levels showed similar tendencies through the differentiation to insulin resistance induction, but the degree of alterations was greater in GLUT4. The mRNA expression levels of insulin resistance markers were remarkably increased during the differentiation, while they were markedly decreased in the insulin resistance model (Figure. 2e-f).

### ***3.3. Effects of cAT-MSC CM treatment on IRS-1 and GLUT4 expressions in the insulin resistance model***

To evaluate the therapeutic effects of cAT-MSC CM on the insulin resistance model, expression levels of IRS-1 and GLUT4 were examined. In the perspective of IRS-1 expression levels, mRNA levels were not changed significantly. However, the protein levels of IRS-1 revealed significant elevation. In GLUT4 expression, both mRNA and protein levels were significantly increased after cAT-MSC CM treatment (Figure. 3).



### ***3.4. Reduced therapeutic effects of cAT-MSC CM with anti-FGF1 neutralizing antibody treatment***

FGF1 in the cAT-MSC CM suspected to play an important role in the alterations in the expression levels of IRS-1 and GLUT4. To verify the hypothesis, an anti-FGF1 neutralizing antibody was added to the cAT-MSC CM treated insulin resistance model. By measuring the protein levels of insulin resistance markers, data confirmed that the improvement of their expression levels was significantly reduced (Figure. 4).

## 4. Discussion

There have been doubts that exogenous insulin supplements for poorly controlled T1DM or T2DM were not enough to maintain the patient in a good condition. In this regard, researches into cell therapy using MSCs have been actively conducted. Mechanisms of their therapeutic effects to DM may be explained through various molecular processes. MSCs have the ability to directly differentiate into IPCs [18, 24, 46] and regulate the immune system to protect pancreatic islet cells from further destruction [1, 8, 17, 22, 51]. In addition, they possess powerful paracrine effects via secretion of a variety of cytokines and growth factors, enabling them to have anti-inflammatory effects and regulate insulin signalling and resulting in the improvement of insulin sensitivity [10, 32, 40, 47]. Several insulin sensitizers (such as biguanides and thiazolidinediones) were considered as therapeutics for DM with insulin resistance. However, long-term medication with these drugs can cause various adverse effects, which highlights the necessity for a new strategy.

In these experiments, 3T3-L1 adipocytes were used to develop an *in vitro* insulin resistance model. There have been diverse studies into inducing insulin resistance in 3T3-L1 cell lines. For example, various agents including TNF- $\alpha$ , Interleukin-1 (IL-1), IL-6, free fatty acids, dexamethasone,

and high insulin were used to make models of insulin resistance in 3T3-L1 adipocytes [20, 30, 35-37, 43]. Lo et al. compared transcriptome analysis data of *in vitro* insulin resistance models and *in vivo* diet-induced obese mouse models. Considering their finding that the TNF- $\alpha$  and hypoxia co-treatment model most closely resembled *in vivo* alterations [29], the model was employed in this experiment. For the purpose of monitoring insulin resistance, IRS-1 and GLUT4 were selected as markers of glucose uptake [41, 49]. IRS-1 is a signalling protein that plays an important role at the initial part of insulin signalling pathway [15]. GLUT4 is a transporter protein that is mainly expressed in the skeletal muscle, adipose tissue, and heart [19]. Its up regulation and translocation to the cell membranes are thought to be critical steps of the insulin responsive increment of glucose uptake [34].

According to early studies, the effectiveness of MSC in DM originated from the potential to trans-differentiation towards IPCs [4, 45]. However, these theories were unable to give a good explanation for the disappointing results of *in vivo* cell tracking studies [5, 27]. Recent studies suggested that the secretory capacity of MSCs would play a crucial role for the therapeutic effects. Gao et al. isolated and injured pancreatic islets *in vitro* and then assessed islet regeneration after the treatment with MSC CM. They found that both  $\beta$  cell replication and islet progenitor differentiation were

promoted and the PI3K / Akt signal pathway was involved [12]. Another study revealed that AT-MSC CM enhanced glucose uptake in 3T3-L1 and C2C12 insulin resistance models [39]. cAT-MSC CM was also utilized to explore the paracrine effect of cAT-MSC in our experiments.

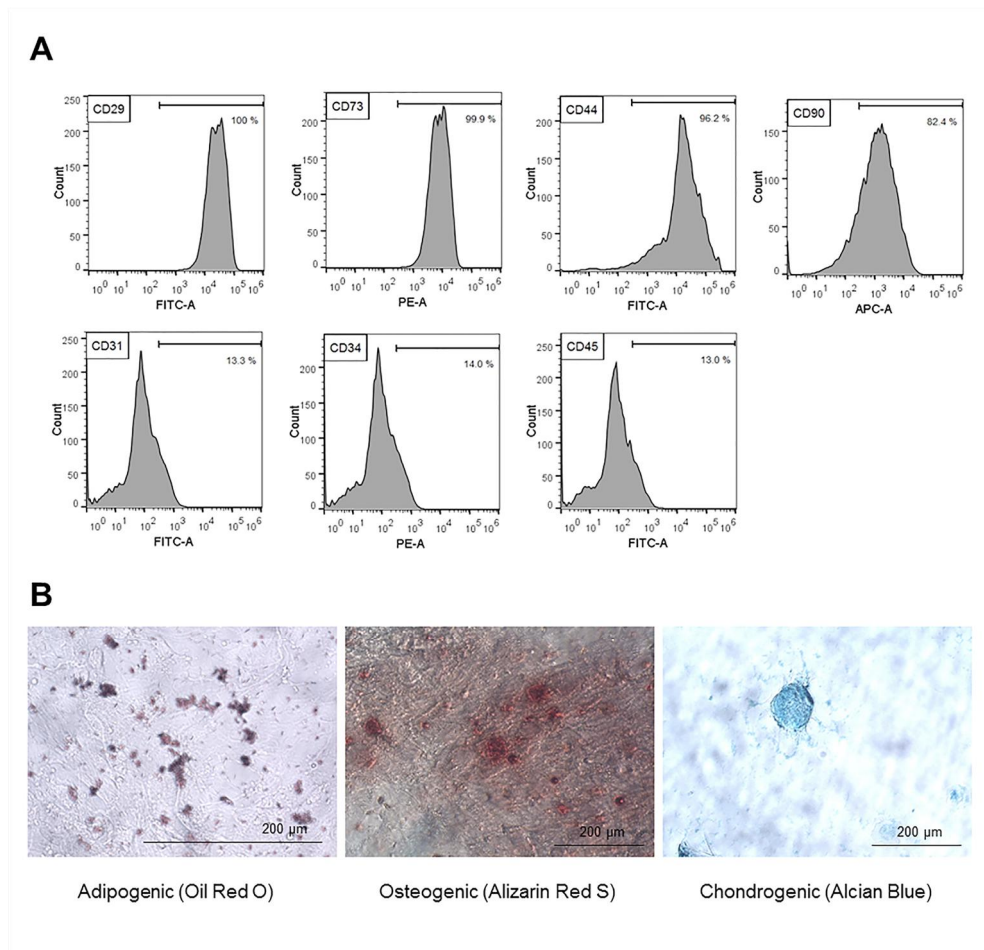
FGF family is a signalling protein group composed of 18 members mediating a variety of biological functions in cell proliferation and developmental processes. Each family member performs unique actions by binding to FGF receptors, which results in activation of intracellular downstream cascades [2]. FGF1, identified as a regulator of adipose tissue remodelling and metabolic homeostasis, is a downstream molecule of peroxisome proliferator-activated receptor- $\gamma$  (PPAR- $\gamma$ ), which is a target of the thiazolidinedione class of insulin sensitizers [21]. It has been established that rFGF1 injection normalized glucose levels and enhanced insulin sensitivity in diabetic mice. What was noteworthy is that the effect was insulin-dependent. Moreover, rFGF1 did not induce side effects that could be incurred by chronic use of traditional insulin sensitizers. Furthermore, genetically modified rFGF1 lacking the mitogen properties of rFGF1 had similar glucose lowering effects to some extent [42], intensifying the possibility of rFGF1 as a substitute of conventional insulin sensitizers. Based on the fact that MSC CM contains considerable amounts of FGF1

[44], I attempted to find out whether FGF1 contributed to the advantageous effects of cAT-MSC CM on insulin resistant models.

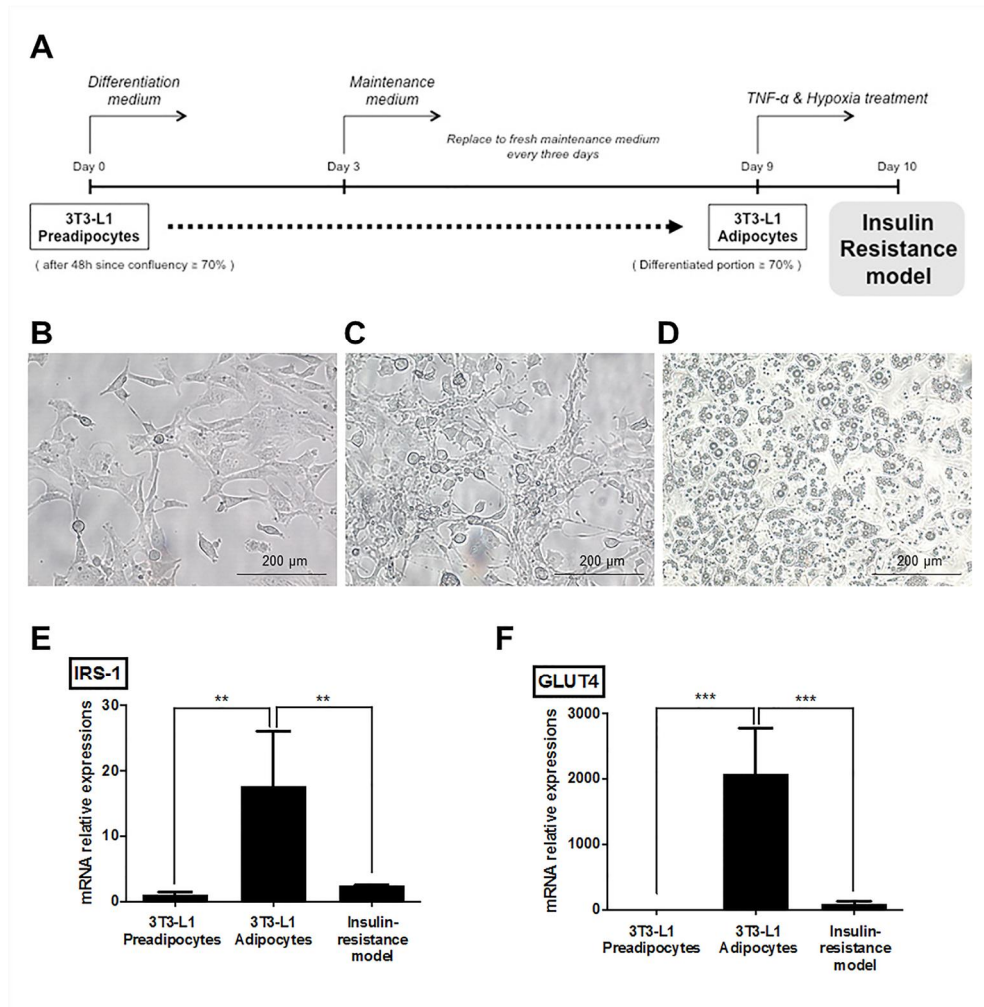
In the present study, gene expression levels of IRS-1 and GLUT4, which were considered as markers of insulin responsive glucose uptake, were evaluated. It was shown that reduced expression levels of these markers in 3T3-L1 insulin resistance models were restored after the treatment of cAT-MSC CM. This finding was consistent with previous research that suggested that MSC could improve insulin resistance through the paracrine signalling of various cytokines and growth factors [39]. Since FGF1 was suspected to have a role in this effect, an anti-FGF1 neutralizing antibody was added into CM-treated 3T3-L1 insulin resistance models and assessed alterations of gene expression levels. The anti-FGF1 antibody group represented diminished insulin resistance marker gene expression levels in comparison to the CM-treated group. This information led to the inference that FGF1 included in cAT-MSC CM is one of the mediators of the signalling pathways that induces therapeutic effects. Although these experiments could not exclude the possibility that various cytokines and growth factors abundantly contained in cAT-MSC CM (other than FGF1) could affect the alterations in expression levels, it is worth mentioning that this study is the first work that revealed FGF1 as a specific mediator of insulin-sensitizing effects of cAT-MSC CM in DM.

## 4. Conclusion

This study identified that the induced insulin resistance in 3T3-L1 cells was ameliorated in the presence of cAT-MSC CM by measuring protein expression levels of IRS-1 and GLUT4. Additional data showed that the improvement was inhibited by the addition of anti-FGF1 neutralizing antibody, suggesting that FGF1 would act as a mediator of the beneficial effects of cAT-MSC CM. It is noteworthy that the present study is the first to reveal a specific functioning component of MSC CM in *in vitro* insulin resistance models, and further studies to find out other effective factors of MSC CM will be necessary.



**Figure 1. Flow cytometry and special staining characterized cAT-MSCs.** (A) Immunophenotypic analysis was performed by flow cytometry using positive markers CD29, CD73, CD44, and CD90 and negative markers CD31, CD34, and CD45. (B) Adipogenic (Oil Red O staining), osteogenic (Alizarin Red S staining), and chondrogenic (Alcian Blue staining) differentiation abilities of cAT-MSCs (from left to right) were confirmed. Original magnification: 400x (left), 200x (middle, right).

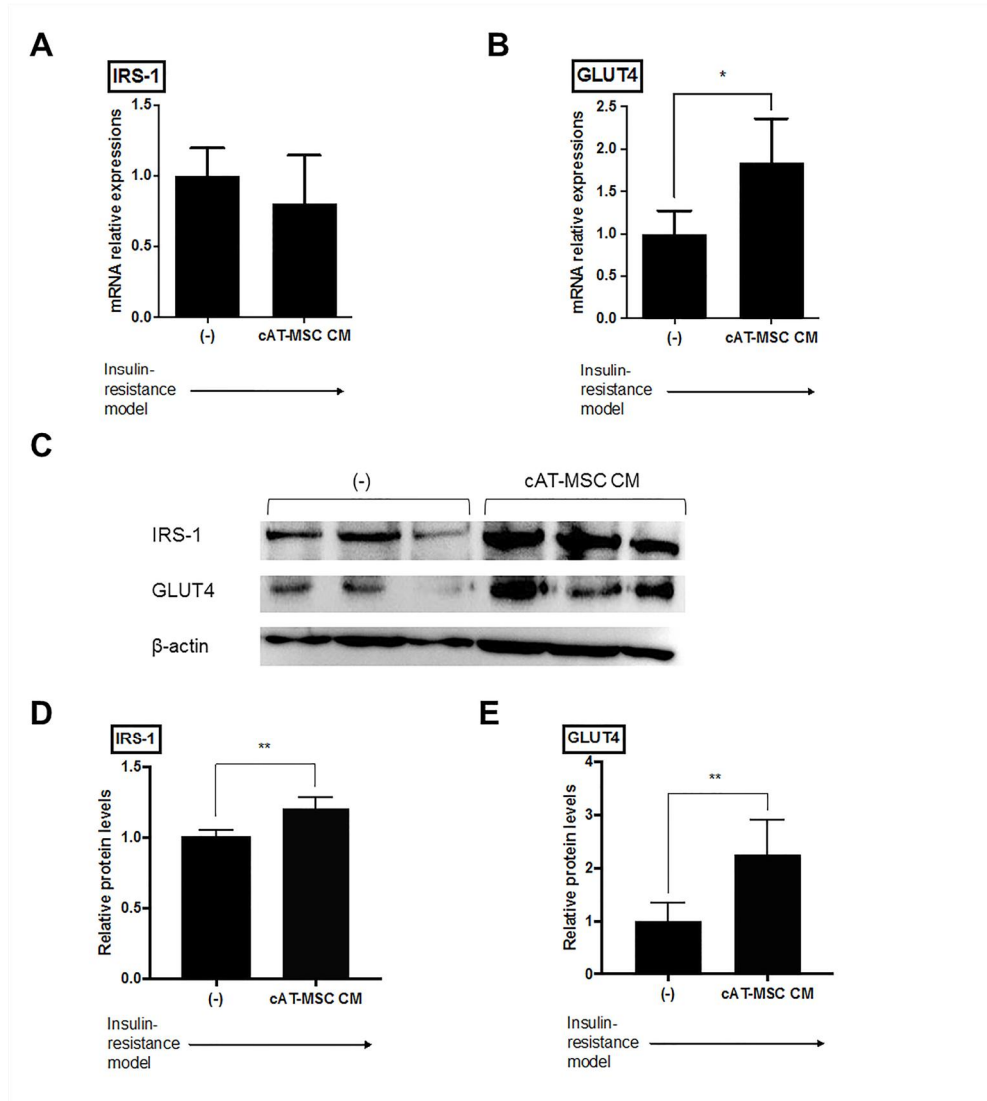


**Figure 2. Insulin resistance was induced in 3T3-L1 adipocytes *in vitro*.**

(A) Adipocyte differentiation of 3T3-L1 cells and following induction of insulin resistance were summarized with the course of time. (B) Fibroblast-like 3T3-L1 preadipocytes at day 0. (C) Differentiating 3T3-L1 cells at day 3. (D) Differentiated 3T3-L1 adipocytes containing cytoplasmic lipid droplets at day 9. Original magnification: 200x. (E, F) Alterations in mRNA

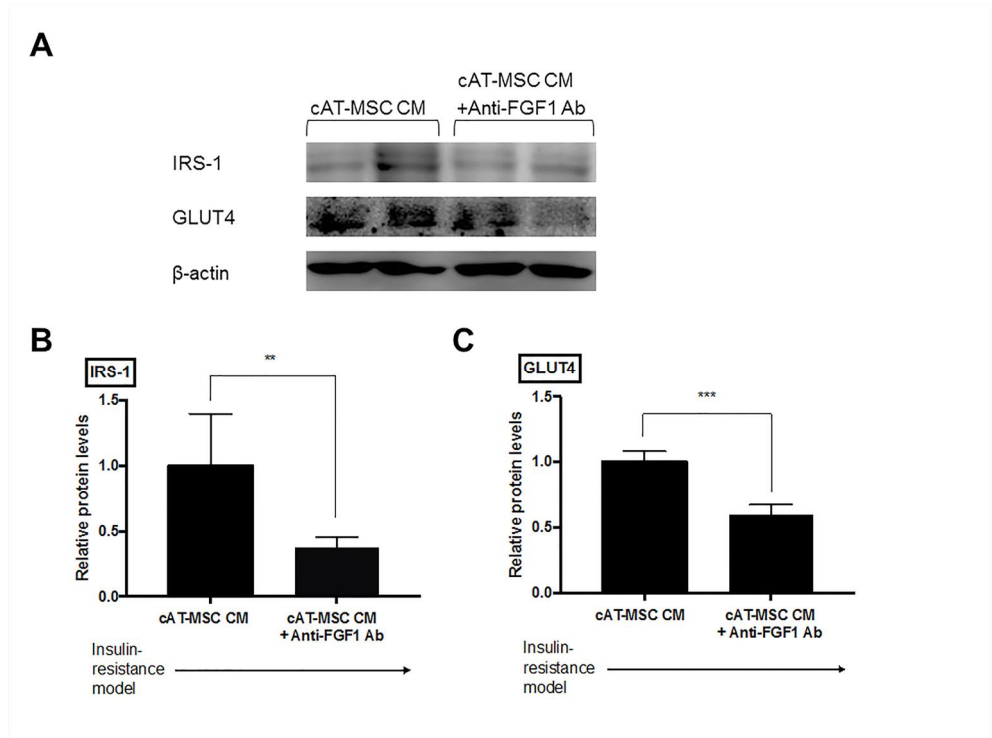


expression levels of IRS-1 and GLUT4 during the differentiation and insulin resistance induction were shown. Data are shown as mean  $\pm$  standard deviation. \*\*P < 0.01, \*\*\*P < 0.001.



**Figure 3. Treatment of cAT-MSC CM affected gene expressions of IRS-1 and GLUT4 in 3T3-L1 insulin resistance models. (A) The mRNA expression levels of IRS-1 were not significantly changed. (B) The mRNA expression levels of GLUT4 showed a significant elevation. (C) Western blot. (-): insulin resistance models of 3T3-L1 cells, cAT-MSC CM: cAT-**

MSC CM-treated insulin resistance models of 3T3-L1 cells. (D-E) Up regulation of IRS-1 and GLUT4 gene expression was observed in CM-treated insulin resistance models. Data are shown as mean  $\pm$  standard deviation. \*P < 0.05, \*\*P < 0.01.



**Figure 4. Gene expressions of IRS-1 and GLUT4 were down regulated after blocking of FGF1 function.** (A) Western blot. (B-C) Relative protein levels of IRS-1 and GLUT4 were lower in the insulin resistance models treated with both cAT-MSC CM and anti-FGF1 antibody than the models treated with cAT-MSC CM alone. Data are shown as mean  $\pm$  standard deviation. \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .

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## 국문 초록

# *In vitro* 인슐린 저항성 모델에 대한 개 지방유래중간엽줄기세포 paracrine 효과의 매개체 fibroblast growth factor-1

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김현진

당뇨 연구 분야에서 중간엽줄기세포를 이용한 세포 치료 연구가 널리 진행되고 있다. 본 연구는 3T3-L1 지방세포에서 유도된 생체 외 인슐린 저항성 모델에서 개 지방유래중간엽줄기세포의 치료 효과를 알아보고, 그 기전을 밝히기 위해 진행되었다. 인슐린 저항성의 지표로서 insulin receptor substrate-1 (IRS-1)과 glucose

transporter type 4(GLUT4)의 유전자 발현 수준을 이용하였다. 개 지방유래중간엽줄기세포를 처리한 인슐린 저항성 모델군은 대조군에 비해 IRS-1와 GLUT4의 상대적인 단백질 발현 수준이 증가되었는데, 이는 당뇨에 대한 개 지방유래중간엽줄기세포의 치료 효과가 분비 인자에 의한 것임을 시사하였다. Fibroblast growth factor-1(FGF1)에 대한 선행연구들을 참고하여 FGF1이 이 기전에 기여할 것이라는 가설을 세우고, anti-FGF1 중화 항체를 줄기세포 처리군에 첨가한 결과, IRS-1과 GLUT4의 단백질 발현 수준이 유의적으로 감소하는 것을 볼 수 있었으며, 이는 앞의 가설을 뒷받침하였다. 따라서 본 연구는 중간엽줄기세포가 인슐린 증감제 대체제를 찾기 위한 연구 대상으로서의 가능성을 지닌다는 점을 확인하였고, paracrine 경로로 작용하는 신호 인자를 새롭게 발견하였다.

**주요어:** fibroblast growth factor-1; 개 지방유래중간엽줄기세포;

당뇨; 인슐린 저항성; 생체 외; 3T3-L1

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